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10	30	50	
CTCGAGCTGGAGTCGGAGTTGTAAACGCTCCACTGACTGATAGAGCGACCGCGCGACCATG			M
70	90	110	
GCGCCCGGAGTGGCCCGCGGCGGACGCCGTACTGGAGGTTGGCGCCTCGGTGGCGCGCGG			
A P G V A R G P T P Y W R L R L G G A A			
130	150	170	
CTGCTCTCTGCTCATCCCGTGGCGCGCGCAGGAGCCTCCCGAGCTGCTTGTCT			
L L L L L I P V A A A Q E P P G A A C S			
190	210	230	
CAGAACACAAACAACTGTGAAGAGTGCTGAAGAAGCTCTCCTGTCCTTTGGTGCAAC			
Q N T N K T C E E C L K N V S C L W C N			
250	270	290	
ACTAACAGGCTTGTCTGGACTACCCAGTTACAAGCGTCTTGGCCACCGGCTTCCCTTG			
T N K A C L D Y P V T S V L P P A F P L			
310	330	350	
TAAATTGAGCTCTGCACGCTGGGAGTTTGTGGTGAACCTTGGAGCGCTGATCATCAC			
370			
CATGTCGGTAG			

FLG. 2

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1  MAPGVARGPTPYWRLRLGGAALLLLIPVAAQEPPGAACSQNTNKTCEE 50
   |||  |||  |||  |||  |||  |||  |||  |||  |||  |||  |||  |||
1  map.....ptr....rllnaallllllmatshqpsgtvarelrcgclk 41

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51 CLKNVSLWCNT...NKACLDYPVTSVLPP 77
    .|.|.|. . . . . . . . .|.|.|.
42 tlpvdfeniqltvtppgphctgteviat 71

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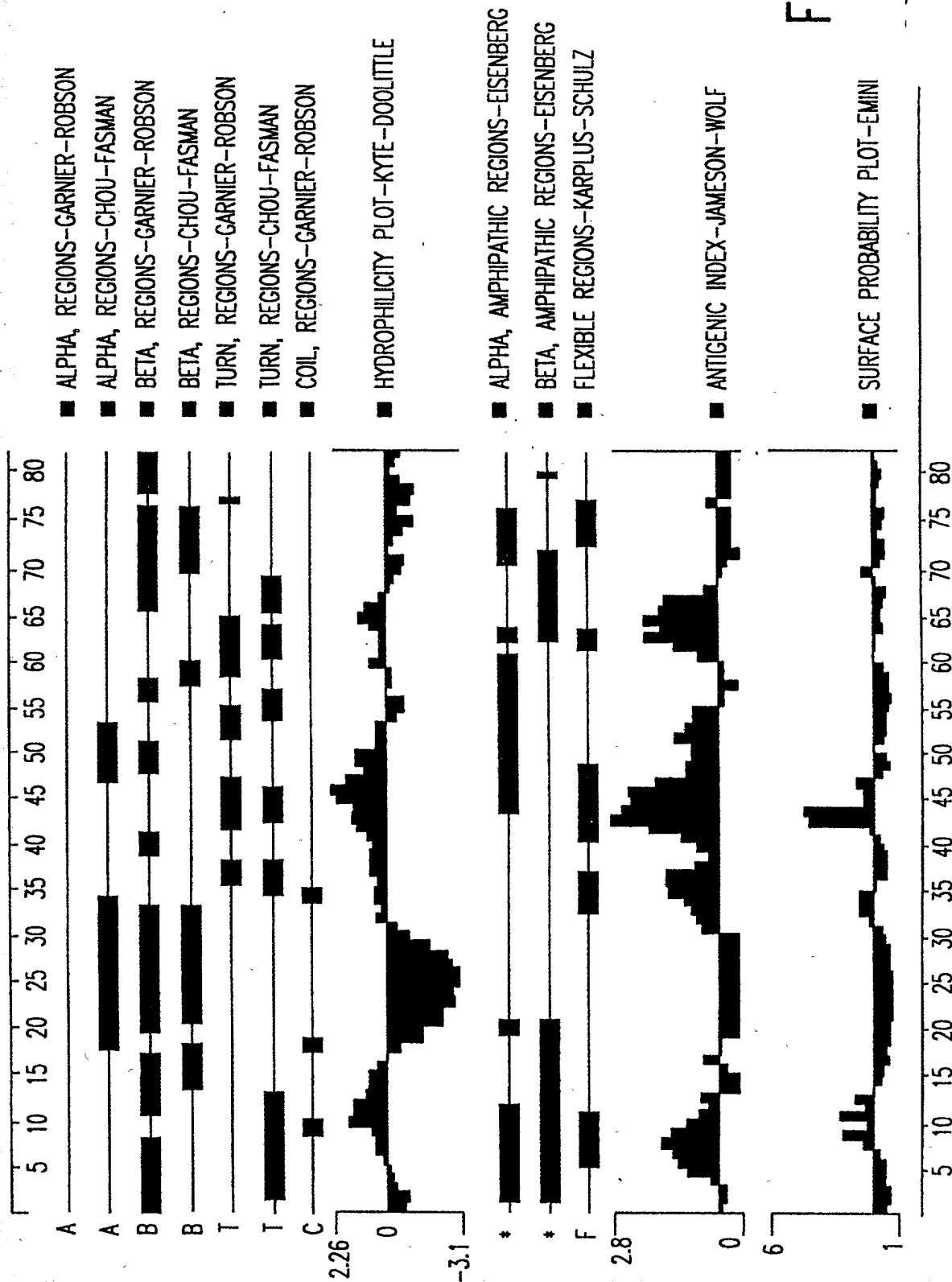


FIG.3